

PCT

## ENTERED

RAW SEQUENCE LISTING

3 <110> APPLICANT: THIERAUCH, KARL-HEINZ

PATENT APPLICATION: US/09/937,100A

DATE: 01/29/2003 P.6

TIME: 15:12:54

Input Set : A:\Merk2309.app

Output Set: N:\CRF4\01292003\I937100A.raw

```
GLIENKE, JENS
      5
             HINZMAN, BERND
             PILARSKY, CHRISTIAN
      8 <120> TITLE OF INVENTION: PROTEIN ISOLATION AND ANALYSIS
     10 <130> FILE REFERENCE: MERCK 2309
     12 <140> CURRENT APPLICATION NUMBER: 09/937,100A
C--> 13 <141> CURRENT FILING DATE: 2002-04-22
     15 <150> PRIOR APPLICATION NUMBER: PCT/GB00/01015
     16 <151> PRIOR FILING DATE: 2000-03-17
     18 <150> PRIOR APPLICATION NUMBER: 9906551.8 GB
     19 <151> PRIOR FILING DATE: 1999-03-23
     21 <150> PRIOR APPLICATION NUMBER: 9907057.5 GB
     22 <151> PRIOR FILING DATE: 1999-03-29
     24 <150> PRIOR APPLICATION NUMBER: 9907641.6 GB
     25 <151> PRIOR FILING DATE: 1999-04-06
     27 <150> PRIOR APPLICATION NUMBER: 9914874.4 GB
     28 <151> PRIOR FILING DATE: 1999-06-28
     30 <150> PRIOR APPLICATION NUMBER: 9915363.7 GB
     31 <151> PRIOR FILING DATE: 1999-07-02
     33 <150> PRIOR APPLICATION NUMBER: 9915677.0 GB
     34 <151> PRIOR FILING DATE: 1999-07-06
    36 <150> PRIOR APPLICATION NUMBER: 9916511.0 GB
     37 <151> PRIOR FILING DATE: 1999-07-14
     39 <150> PRIOR APPLICATION NUMBER: 9920503.1 GB
     40 <151> PRIOR FILING DATE: 1999-08-31
     42 <150> PRIOR APPLICATION NUMBER: 9922285.3 GB
     43 <151> PRIOR FILING DATE: 1999-09-21
     45 <160> NUMBER OF SEQ ID NOS: 69
     47 <170> SOFTWARE: PatentIn Ver. 2.1
     49 <210> SEQ ID NO: 1
     50 <211> LENGTH: 24
    51 <212> TYPE: DNA
     52 <213> ORGANISM: Artificial Sequence
     54 <220> FEATURE:
    55 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
             oligonucleotide for an 8 amino acid barcode peptide
     58 <220> FEATURE:
     59 <221> NAME/KEY: CDS
     60 <222> LOCATION: (1)..(24)
     61 <223> OTHER INFORMATION:
     63 <220> FEATURE:
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64 <221> NAME/KEY: misc feature

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```
Input Set : A:\Merk2309.app
                     Output Set: N:\CRF4\01292003\I937100A.raw
     65 <222> LOCATION: (1)..(1)
     66 <223> OTHER INFORMATION: n=a,t,g,c
W--> 68 <220>
     69 <221> NAME/KEY: misc feature
     70 <222> LOCATION: (4)..(4)
     71 <223> OTHER INFORMATION: n=a,t,g,c
     73 <220> FEATURE:
     74 <221> NAME/KEY: misc feature
     75 <222> LOCATION: (7)...(7)
     76 <223> OTHER INFORMATION: n=a,t,g,c
     78 <220> FEATURE:
     79 <221> NAME/KEY: misc feature
     80 <222> LOCATION: (10)..(10)
     81 <223> OTHER INFORMATION: n=a,t,q,c
     83 <220> FEATURE:
     84 <221> NAME/KEY: misc feature
     85 <222> LOCATION: (14)..(14)
     86 <223> OTHER INFORMATION: k=q,t
     88 <220> FEATURE:
     89 <221> NAME/KEY: misc feature
     90 <222> LOCATION: (16)..(16)
     91 <223> OTHER INFORMATION: v=a,q,c
     93 <220> FEATURE:
     94 <221> NAME/KEY: misc feature
     95 <222> LOCATION: (20)..(20)
     96 <223> OTHER INFORMATION: n=a,q,t,c
     98 <220> FEATURE:
     99 <221> NAME/KEY: misc feature
     100 <222> LOCATION: (21)..(21)
     101 <223> OTHER INFORMATION: v=a,q,c
     103 <220> FEATURE:
     104 <221> NAME/KEY: misc feature
     105 <222> LOCATION: (23)..(23)
     106 <223> OTHER INFORMATION: n=a,t,g,c
     108 <400> SEQUENCE: 1
                                                                                 24
W--> 109 nac ncc ngg ntg tkc vag gnv cnt
W--> 110 Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     111 1
     114 <210> SEQ ID NO: 2
     115 <211> LENGTH: 8
     116 <212> TYPE: PRT
     117 <213> ORGANISM: Artificial Sequence
     119 <220> FEATURE:
     120 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     121
               barcode peptide
     123 <220> FEATURE:
     124 <221> NAME/KEY: misc feature
     125 <222> LOCATION: (1)..(1)
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126 <223> OTHER INFORMATION: The 'Xaa' at location 1 stands for Asn, Asp, His, or Tyr.

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Input Set : A:\Merk2309.app Output Set: N:\CRF4\01292003\I937100A.raw 128 <220> FEATURE: 129 <221> NAME/KEY: misc feature 130 <222> LOCATION: (2)..(2) 131 <223> OTHER INFORMATION: The 'Xaa' at location 2 stands for Thr, Ala, Pro, or Ser. 133 <220> FEATURE: 134 <221> NAME/KEY: misc feature 135 <222> LOCATION: (3)..(3) 136 <223> OTHER INFORMATION: The 'Xaa' at location 3 stands for Arg, Gly, or Trp. 138 <220> FEATURE: 139 <221> NAME/KEY: misc feature 140 <222> LOCATION: (4)..(4) 141 <223> OTHER INFORMATION: The 'Xaa' at location 4 stands for Met, Val, or Leu. 143 <220> FEATURE: 144 <221> NAME/KEY: misc feature 145 <222> LOCATION: (5)..(5) 146 <223> OTHER INFORMATION: The 'Xaa' at location 5 stands for Cys, or Phe. 148 <220> FEATURE: 149 <221> NAME/KEY: misc feature 150 <222> LOCATION: (6)..(6) 151 <223> OTHER INFORMATION: The 'Xaa' at location 6 stands for Lys, Glu, or Gln. 153 <220> FEATURE: 154 <221> NAME/KEY: misc feature 155 <222> LOCATION: (7)..(7) 156 <223> OTHER INFORMATION: The 'Xaa' at location 7 stands for Glu, Asp, Gly, Ala, or 158 <220> FEATURE: 159 <221> NAME/KEY: misc feature 160 <222> LOCATION: (8)..(8) 161 <223> OTHER INFORMATION: The 'Xaa' at location 8 stands for His, Arg, Pro, or Leu. 163 <400> SEOUENCE: 2 W--> 164 Xaa Xaa Xaa Xaa Xaa Xaa Xaa 165 1 168 <210> SEQ ID NO: 3 169 <211> LENGTH: 14 170 <212> TYPE: PRT 171 <213> ORGANISM: Artificial Sequence 173 <220> FEATURE: 174 <223> OTHER INFORMATION: Description of Artificial Sequence: Linker peptide 177 <400> SEQUENCE: 3 178 Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Val Asp 179 1 182 <210> SEQ ID NO: 4 183 <211> LENGTH: 8 184 <212> TYPE: PRT 185 <213> ORGANISM: Artificial Sequence 187 <220> FEATURE: 188 <223> OTHER INFORMATION: Description of Artificial Sequence: Flag 189 epitope peptide 191 <400> SEQUENCE: 4

RAW SECUENCE LISTING

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Val.

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Input Set : A:\Merk2309.app

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```
192 Met Asp Tyr Lys Asp Asp Asp Lys
193 1
196 <210> SEQ ID NO: 5
197 <211> LENGTH: 53
198 <212> TYPE: DNA
199 <213> ORGANISM: Artificial Sequence
201 <220> FEATURE:
202 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
          RD5' Flag
205 <400> SEQUENCE: 5
                                                                           53
206 gcggatccca tatggactac aaagacgatg acgacaaaca ggtgcagctg cag
209 <210> SEQ ID NO: 6
210 <211> LENGTH: 35
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
216
       RD3'
218 <400> SEQUENCE: 6
                                                                           35
219 gcgaattcgt ggtggtggtg gtggtgtgac tctcc
222 <210> SEO ID NO: 7
223 <211> LENGTH: 50
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
229
          Fos1for
231 <400> SEQUENCE: 7
                                                                           50
232 atggaattcc tcgagaccga caccctacag gcggaaaccg accagctgga
235 <210> SEO ID NO: 8
236 <211> LENGTH: 50
237 <212> TYPE: DNA
238 <213> ORGANISM: Artificial Sequence
240 <220> FEATURE:
241 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
242
       Fos80rev
244 <400> SEQUENCE: 8
245 tcgcgatttc ggtttgcagc gcggattttt cgtcttccag ctggtcggtt
                                                                           50
248 <210> SEQ ID NO: 9
249 <211> LENGTH: 50
250 <212> TYPE: DNA
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
255
         Fos71for
257 <400> SEQUENCE: 9
258 aaaccqaaat cgcgaacctg ctgaaagaaa aagaaaagct ggagttcatc
                                                                           50
261 <210> SEO ID NO: 10
262 <211> LENGTH: 50
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Input Set : A:\Merk2309.app

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263 <212> TYPE: DNA
264 <213> ORGANISM: Artificial Sequence
266 <220> FEATURE:
267 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
          Fos155rev
270 <400> SEQUENCE: 10
                                                                            50
271 ggaagettga atteegeegg acggtgtgee geeaggatga acteeagett
274 <210> SEO ID NO: 11
275 <211> LENGTH: 18
276 <212> TYPE: DNA
277 <213> ORGANISM: Artificial Sequence
279 <220> FEATURE:
280 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
281
         Fosl fS
283 <400> SEOUENCE: 11
                                                                            18
284 atggaattcc tcgagacc
287 <210> SEQ ID NO: 12
288 <211> LENGTH: 18
289 <212> TYPE: DNA
290 <213> ORGANISM: Artificial Sequence
292 <220> FEATURE:
293 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
          Fos155 rS
294
296 <400> SEOUENCE: 12
                                                                            18
297 ggaagcttga attccgcc
300 <210> SEQ ID NO: 13
301 <211> LENGTH: 28
302 <212> TYPE: DNA
303 <213> ORGANISM: Artificial Sequence
305 <220> FEATURE:
306 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer I
          for 340 VH amplification
309 <400> SEOUENCE: 13
                                                                            28
310 cagctgcagg agtctggggg aggcttag
313 <210> SEQ ID NO: 14
314 <211> LENGTH: 36
315 <212> TYPE: DNA
316 <213> ORGANISM: Artificial Sequence
318 <220> FEATURE:
319 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer II
          for 340 VH amplification
320
322 <400> SEQUENCE: 14
                                                                            36
323 teagtagacg gtgaccgagg tteettgace ceagta
326 <210> SEQ ID NO: 15
327 <211> LENGTH: 26
328 <212> TYPE: DNA
329 <213> ORGANISM: Artificial Sequence
331 <220> FEATURE:
332 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer I
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/937,100A

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## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:1; N Pos. 1,4,7,10,20,23
Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:2; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:27; N Pos. 21,24
Seg#:28; N Pos. 21,24
Seq#:29; N Pos. 21,24
Seq#:30; N Pos. 21,24
Seq#:31; N Pos. 21,24
Seq#:32; N Pos. 7,10
Seq#:33; N Pos. 7,10
Seq#:34; N Pos. 7,10
Seq#:35; N Pos. 7,10
Seq#:36; N Pos. 7,10
Seg#:58; N Pos. 31,32,34,35,37,38,40,41,43,44,46,47,49,50
Seq#:58; Xaa Pos. 11,12,13,14,15,16,17
Seg#:59; Xaa Pos. 11,12,13,14,15,16,17
Seq#:60; N Pos. 28,29,31,32,34,35,37,38,40,41,43,44,46,47
Seq#:61; N Pos. 14,15,17,18,20,21,23,24,26,27,29,30,32,33,35,36,38,39,41,42
Seg#:61; Xaa Pos. 5,6,7,8,9,10,11,12,13,14
Seq#:62; Xaa Pos. 5,6,7,8,9,10,11,12,13,14
Seq#:63; N Pos. 29,30,32,33,35,36,38,39,41,42,44,45,47,48,50,51,53,54,56,57
Seg#:64; N Pos. 15,18,19,22,25,28,38,41
Seg#:64; Xaa Pos. 7,8,9,10,11,12,13,14
Seq#:65; Xaa Pos. 7,8,9,10,11,12,13,14
Seg#:66; N Pos. 14,17,27,30,33,36
Seq#:67; N Pos. 15,18,19,22,25,28,38,41,48,51,52,55,58,61,71,74
Seq#:67; Xaa Pos. 7,8,9,10,11,12,13,14,18,19,20,21,22,23,24,25
Seq#:68; Xaa Pos. 7,8,9,10,11,12,13,14,18,19,20,21,22,23,24,25
```

Seg#:69; N Pos. 14,17,27,30,33,36,47,50,60,63,66,69

## VERIFICATION SUMMARY

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Input Set : A:\Merk2309.app

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```
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:68 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:61
L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:24 L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:525 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:617 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:640 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
L:663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:709 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:1017 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:58, Line#:1010
L:1083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:0
L:1084 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:48
L:1087 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:48
L:1088 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:47
L:1151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
L:1154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:16
L:1239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:0
L:1263 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:61, Line#:1256
L:1359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0
L:1360 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:49
L:1448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0
L:1566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:0
L:1589 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:64,Line#:1582
L:1645 \text{ M}:341 \text{ W}: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0
L:1646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:48
L:1703 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:0
L:1762 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:0
L:1784 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:67,Line#:1777
L:1890 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:0
L:1891 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:48
L:1894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:48
L:1895 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:87
L:1989 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:0
L:1992 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:16
L:2098 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:0
L:2099 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:60
```